## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/553,298
Source:	PG/10.
Date Processed by STIC:	10/25/05

## ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 10/25/2005

PATENT APPLICATION: US/10/553,298

TIME: 10:41:40

Input Set : A:\sequence.txt

Output Set: N:\CRF4\10252005\J553298.raw

3 <110> APPLICANT: NeuroNova AG

5 <120> TITLE OF INVENTION: A Method for Diagnosing and Treating Affective Disorders

7 <130> FILE REFERENCE: XXX

C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/553,298

C--> 9 <141> CURRENT FILING DATE: 2005-10-14

9 <160> NUMBER OF SEQ ID NOS: 111

11 <170> SOFTWARE: PatentIn version 3.1

## **ERRORED SEQUENCES**

4643 <210> SEQ ID NO: 111

4644 <211> LENGTH: 20

4645 <212> TYPE: DNA

4646 <213> ORGANISM: Homo sapiens

4648 <400> SEQUENCE: 111

4649 cacagagett tgcaggtgaa

E--> 4653/1 E--> 4656 1

Does Not Comply orrected Diskette Neede

20

VERIFICATION SUMMARY DATE: 10/25/2005 PATENT APPLICATION: US/10/553,298 TIME: 10:41:41

Input Set : A:\sequence.txt

Output Set: N:\CRF4\10252005\J553298.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:19 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:24 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:28 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:21 L:29 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:33 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:26 L:34 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:38 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:31 L:39 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:43 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:36 L:44 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:48 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:41 L:49 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:53 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:46 L:54 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:58 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:51 L:59 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:63 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:56 L:64 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:68 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:61 L:69 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:73 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:66 L:74 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:78 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:71 L:79 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1 L:83 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:76 L:83 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:81 L:1981 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:2,Line#:1979 L:2328 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:2320 L:2339 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:2331 L:4653 M:254 E: No. of Bases conflict, this line has no nucleotides. M:254 Repeated in SeqNo=111